

Fig. 1**SEQ ID NO: 1**

GGGCCGGGCAGGGCCGGGCGTGGGCCGGCAGGAAGATGGCGAACGTG
GGGCTGCAGTTCCAGGCGAGCGCGGGGACTCGGACCCACAGAGCCGG
CCCCTGCTGCTGCTCGGGCAGCTGCACCACCTGCACCGCGTGCCTGG
AGCCACGTCCGGGAAGCTGCAGCCCCGGTCACCGAGGAGCTCTGG
CAGGCTGCCCTGAGCACGCTCAACCCCAACCCACGGACAGCTGTCCC
CTCTACCTAACTACGCCACCGTGGCTGCCCTGCCCTGCAGGGTGAGCC
GGCACAAACAGCCCCTGGCCGCCACTTCATCACGCGCTGGTGCAGGA
CCTGCCTGCCGCCGGAGCGCATCGCTGCATTGTGATGGTCTGCGAGC
AGCCAGAGGTCTTGCTTCCGCCTGTGCCCTGGCCCGGGCCTCCCGC
TGTTCACCCACCGCTCAGGTGCCTCTCGCGCTTGGAGAAGAACAGG
TCACCGTGGAGTTTCTGGTGGACAAGACAACGGGCCGGTGGAGG
TGTCCACATTGCAGTGCTTAGCGAATGCCACAGACGGCGTGCAGCTAG
CAGCCCGCATCGTGGACACACCCCTGCAATGAGATGAACACCGACACCT
TCCTCGAGGAGATTAACAAAGCTGGAAAGGAGCTGGGATCATCCCAA
CCATCATCCGGATGAGGAACCTGAAGACGAGAGGATTGGAGGAATCT
ATGGGGTTGGCAAAGCCGCCCTGCATCCCCCAGCCCTGGCGTCCTCA
GCCACACCCAGATGGAGCCACGCAGACCATCGCCTGGTGGCAAAG
GCATCGTCTATGACACTGGAGGCCTCAGCATCAAAGGAAGACTACCA
TGCCGGGATGAAGCGAGACTGCGGGGTGCTGCGGCCGTCTGGGG
CCTTCAGAGCCGAATCAAGCAGGGTTCAAAGACAACCTCCACGCTG
TGTTCTGCTTGGCTGAGAACTCGTGGGCCAATGCGACAGGGCCAG
ATGACATCCACCTGCTGTACTCAGGGAAAGACGGTGGAAATCAACAACA
CGGATGCCGAGGGCAGGCTGGTGCAGATGGCGTGCCTATGCTT
GCAAGGACCTGGGGCCGACATCATCCTGGACATGCCACCCCTGACCG
GGGCTCAGGGCATTGCCACAGGGAAAGTACCAAGCCCGGTGCTCACCA
ACAGCGCTGAGTGGAGGCCCTGTGTGAAGGCGGGCAGGAAGTGTG
GGGACCTGGTGCACCCGCTGGTCTACTGCCCGAGCTGCACCTCAGCG
AGTTCACCTCAGCTGTGGCGGACATGAAGAACTCAGTGGCGGACCGAG

ACAAACAGCCCCAGCTCCTGTGCTGGCCTCTTCATCGCCTCACACATCG
GCTTCGACTGGCCCGGAGTCTGGGTCCACCTGGACATTGCTGCACC
TGCATGCTGGTGAGCGAGCCACAGGCTTCGGTGTGGCCCTCTGCTGG
CGCTCTTCGGCCGTGCCTCTGAGGACCCCTGCTGAACCTGGTGTCCC
CACTGGGCTGTGAGGTGGATGTCGAGGGAGGGGACCTGGGAGGGACT
CCAAGAGACGCAGGCTTGTGTGAGCCTCCTGCCTCGGCCCTGACAAAC
GGGGATCTTTACCTCACTTGCACTGATTAATTTAAGCAATTGAAA
GATTGCCCTTC

Fig. 2**SEQ ID NO: 2**

MVCEQPEVFASACALARAFPLFTHRSGASRRLEKKTVTVEFFLVGQDN
GPVEVSTLQCLANATDGVRЛАARIVDTPCNEMNTDTFLEEINKVGKEL
GIIPTIIRDEELKTRGFГGIYVGVKAALHPPALAVLSHTPDGATQTIA
WVGKGIVYDTGGLSIKGKTTMPGMKRDCGGAAVLGAFRAAIKQGFKD
NLHAVFCLAENSVPNATRPDDIHLLYSGKTVEINNTDAEGLVLA
VSYACKDLGADIILDMATLTGAQGIATGKYHAAVLTNSAEWEAACVKA
GRKCGDLVHPLVYCPHELHFSEFTSAVADMKN
VADRDNSPSSCAGLFI
ASHIGFDWPGVWVHLDIAAPVHAGERATGFGVALLLALFGRASEDPLL
NLVSPLGCEVDVEEGDLGRDSKRRRLV

Fig. 3**SEQ ID NO: 3**

5' - TTTCTGGTGGACAAGACA - 3'

Fig. 4**SEQ ID NO: 4**

5' - CATTGCTAAGCACTGCAATG - 3'

Fig. 5

SEQ ID NO: 5

5 ' - CGGGCCGGTGGAGGTGTC - 3 '